Application No.: 10/646,985 Docket No.: COTH-P03-504

Amendments to the Claims

Please amend claim 1. This listing of claims will replace all prior versions and listings of claims in the application:

Listing of Claims:

- 1. (Currently amended) A method for removing the 3'-untranslated region of a population of DNA molecules, wherein each DNA molecule in said population of DNA molecules comprises an open reading frame and a 3'-untranslated region, said method comprising:
- (a) providing a population of DNA molecules, each of said DNA molecules terminating in a 3' overhang upstream of the open reading frame and in a blunt end downstream of the open reading frame, wherein each of said DNA molecules is double-stranded; and
- (b) treating each of said DNA molecules first with an exonuclease III and then with a single-stranded nuclease under conditions that allow removal of said 3'-untranslated region,

wherein the sequential treatment of said DNA molecules with the exonuclease III and the nuclease specifically removes the 3'-untranslated region from the end downstream of the open reading frame but not the open reading frame.

- 2. (Canceled)
- 3. (Original) The method of claim 1, wherein said nuclease is Mung bean nuclease.
- 4. (Original) The method of claim 1, wherein step (b) further results in removal of the stop codon of said open reading frame.
- 5. (Original) The method of claim 1, wherein each of said DNA molecules is a cDNA produced by reverse transcription from an mRNA sequence.
- 6. (Original) The method of claim 1, wherein said population comprises at least 10 DNA molecules.

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7. (Original) The method of claim 1, wherein said population comprises at least 10² DNA molecules.

- 8. (Original) The method of claim 1, wherein said population comprises at least 10³ DNA molecules.
- 9. (Original) The method of claim 1, wherein said population comprises at least 10⁴ DNA molecules.
- 10. (Original) The method of claim 1, wherein said population comprises at least 10⁵ DNA molecules.
- 11. (Original) The method of claim 1, wherein said population comprises at least 10⁶ DNA molecules.